

| | | | | | |
|------|------------|------------|------------|------------|------|
| -3 | TTCATGGCAG | CTCAGCACAT | CATCTCCATG | GCTGCCATGG | 37 |
| 38 | CATCCCTCCT | TGTAGTACTC | TCGGCAATCC | CGAGAGGCGT | 77 |
| 78 | GGAATCCATT | GGGGTCTGCA | ATGGAATGGA | CGGTGACAAC | 117 |
| 118 | CTCCCCCAGC | CCGCCGACGT | CGTCAACCTC | TACAAGTCCA | 157 |
| 158 | ACAACATAGC | TGGCATGCGA | CTCTACAGCC | CCGACCAAGC | 197 |
| 198 | CACTCTCCAG | GCCCTCCAGG | GCTCTAACAT | CTACCTCATC | 237 |
| 238 | CTCGACGTCC | CCAACTCCGA | CCTCCAAAAC | ATTGCCTCCG | 277 |
| 278 | ACCAATCCGC | CGCCACCAAC | TGGGTCCAAA | CCAACGTCCA | 317 |
| 318 | AGCCTACCCA | AACGTTGCCT | TCCGATACAT | CGCCGTCGGA | 357 |
| 358 | AACGAAGTCA | TCCCCGGCGG | CCAAGCTCAG | TACGTCCTCC | 397 |
| 398 | CAGCCATGAA | CAACATACAG | TCCGCCCTCT | CCTCTGCCGG | 437 |
| 438 | CCTTCAGAAC | ATCAAGGTCT | CCACATCAGT | CTCCTTCGGC | 477 |
| 478 | GTCGTCGGTA | CCTCATATCC | CCCCTCAGGT | GGCTCCTTCT | 517 |
| 518 | CTTCCGATGC | ATCGTCGACA | TTGGGTCCAA | TCATACAGTT | 557 |
| 558 | TCTAGCCAGC | AATGGCTCCC | CATTACTTGC | CAACATCTAC | 597 |
| 598 | CCCTACTTGA | GCTATGCTGG | CAACTCCGGA | TCCATCGACC | 637 |
| 638 | TCTCATACGC | CCTCTTTACT | GCATCTGGTA | CAGTCGTACA | 677 |
| 678 | GGACGGGTCC | TACGCTTACA | ACAACCTCTT | CGATGCCATG | 717 |
| 718 | GTCGACGCAT | TGTACTCGGC | CCTGGAGAGC | GCCGGAGGGC | 757 |
| 758 | CGAATGTCCC | TGTTGTCTGT | TCGGAGAGTG | GCTGGCCGTC | 797 |
| 798 | AGCGGGCGGG | ACAGCGGCGA | CGGTGTCTAA | TGCGCAGACT | 837 |
| 838 | TACAATTCCA | ATTTGATCAA | CCATGTGGGT | CAGGGGACGC | 877 |
| 878 | CGAAGAGGCC | AGGGGCGATT | GAGACCTACA | TATTTGCCAT | 917 |
| 918 | GTTCAACGAG | GATCAGAAGC | AGCCGCAAGG | GATTGAGAAT | 957 |
| 958 | AACTTTGGGC | TGTTTTACCC | TAACGAACAG | CCTGTCTATT | 997 |
| 998 | CGATCAGCTT | CACCTGAGAA | ATTTGATCAG | ATGAAATATA | 1037 |
| 1038 | AAATAAAGGT | CTTATATTGT | AAGGCAAAGC | TCGTAATTGA | 1077 |
| 1078 | TAGCCATCTA | GTAATATAGC | TCCGGCTAAT | TAAAACTATA | 1117 |
| 1118 | AAATA | | | | 1122 |

Fig. 4. The nucleotide sequence of the full-length cDNA of *LP_{Glu}I*

(incorporating SEQ ID NO: 2).

1 MAAQHIISMA AMASLLVVLS AIPRGVESIG VCNGMDGDNL PQPADVVNLY
51 KSNNIAGMRL YSPDQATLQA LQGSNIYLIL DVPNSDLQNI ASDQSAATNW
101 VQTNVQAYPN VAFRYIAVGN EVIPGGQAQY VLPAMNNIQS ALSSAGLQNI
151 KVSTSVSFGV VGTSYPPSAG SFSSDASSTL GPIIQFLASN GSPLLANIYP
201 YLSYAGNSGS IDLSYALFTA SGTVVQDGSY AYNNLFDAMV DALYSALESA
251 GGPNVVPPVVS ESGWPSAGGT AATVSNAQTY NSNLINHVQG GTPKRPGAIE
301 TYIFAMFNED QKQPQGIENN FGLFYPNEQP VYSISFT

Fig. 5. The putative amino acid sequence of LPGlu1 (SEQ ID NO: 1).